

SEQLIST.TXT

SEQUENCE LISTING

<110> Albert, Lai

<120> NOVEL SPLICE VARIANTS OF HUMAN Dkk11

<130> PP023359.0003

<140> 10/574182

<141> 2007-05-31

<150> PCT/US04/34256

<151> 2004-09-30

<150> 60/507682

<151> 2003-09-30

<160> 28

<170> FastSEQ for Windows Version 4.0

<210> 1

<211> 819

<212> DNA

<213> Homo sapien

<400> 1

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cgggatggac tccgcaaggg gacccacaag gacgtcctag aagaggggac cgagagctcc 660
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<210> 2

<211> 242

<212> PRT

<213> Homo sapien

<400> 2

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Met Gly Glu Ala Ser Pro Pro Ala Pro Ala Arg Arg His Leu Leu Val
1      5      10      15
Leu Leu Leu Leu Ser Thr Leu Val Ile Pro Ser Ala Ala Ala Pro
20     25     30
Ile His Asp Ala Asp Ala Gln Glu Ser Ser Leu Gly Leu Thr Gly Leu
35     40     45
Gln Ser Leu Leu Gln Gly Phe Ser Arg Leu Phe Leu Lys Gly Asn Leu
50     55     60
Leu Arg Gly Ile Asp Ser Leu Phe Ser Ala Pro Met Asp Phe Arg Gly
65     70     75     80
Leu Pro Gly Asn Tyr His Lys Glu Glu Asn Gln Glu His Gln Leu Gly
85     90     95
Asn Asn Thr Leu Ser Ser His Leu Gln Ile Asp Lys Arg Thr Asp Asn

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      100      105      110
Lys Thr Gly Glu Val Leu Ile Ser Glu Asn Val Val Ala Ser Ile Gln
      115      120      125
Pro Ala Glu Gly Ser Phe Glu Gly Asp Leu Lys Val Pro Arg Met Glu
      130      135      140
Glu Lys Glu Ala Leu Val Pro Ile Gln Lys Ala Thr Asp Ser Phe His
      145      150      155      160
Thr Glu Leu His Pro Arg Val Ala Phe Trp Ile Ile Lys Leu Pro Arg
      165      170      175
Arg Arg Ser His Gln Asp Ala Leu Glu Gly Gly His Trp Leu Ser Glu
      180      185      190
Lys Arg His Arg Leu Gln Ala Ile Arg Asp Gly Leu Arg Lys Gly Thr
      195      200      205
His Lys Asp Val Leu Glu Glu Gly Thr Glu Ser Ser Ser His Ser Arg
      210      215      220
Leu Ser Pro Arg Lys Thr His Leu Leu Tyr Ile Leu Arg Pro Ser Arg
      225      230      235      240
Gln Leu

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<210> 3
 <211> 733
 <212> DNA
 <213> Homo sapien

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agagagctcc ttgggtctca caggcctcca gagcctactc caaggcttca gccgactttt 180
cctgaaaggt aacctgcttc ggggcataga cagcttattc tctgccccca tggacttccg 240
gggcctccct gggaactacc acaaagagga gaaccaggag caccagctgg ggaacaacac 300
cctctccagc cacctccaga tcgacaagat gaccgacaac aagacaggag aggtgctgat 360
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ccaccaggat gccctggagg gcggccactg gctcagcgag aagcgacacc gcctgcaggc 600
catccgggat ggactccgca aggggaccca caaggacgtc ctagaagagg ggaccgagag 660
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tcggcagctg tag 733

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<210> 4
 <211> 242
 <212> PRT
 <213> Homo sapien

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<400> 4
Met Gly Glu Ala Ser Pro Pro Ala Pro Ala Arg Arg His Leu Leu Val
1      5      10      15
Leu Leu Leu Leu Leu Ser Thr Leu Val Ile Pro Ser Ala Ala Ala Pro
20     25     30
Ile His Asp Ala Asp Ala Gln Glu Ser Ser Leu Gly Leu Thr Gly Leu
35     40     45
Gln Ser Leu Leu Gln Gly Phe Ser Arg Leu Phe Leu Lys Gly Asn Leu
50     55     60
Leu Arg Gly Ile Asp Ser Leu Phe Ser Ala Pro Met Asp Phe Arg Gly
65     70     75     80
Leu Pro Gly Asn Tyr His Lys Glu Glu Asn Gln Glu His Gln Leu Gly
85     90     95
Asn Asn Thr Leu Ser Ser His Leu Gln Ile Asp Lys Met Thr Asp Asn
100    105    110
Lys Thr Gly Glu Val Leu Ile Ser Glu Asn Val Val Ala Ser Ile Gln
115    120    125

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Pro Ala Glu Gly Ser Phe Glu Gly Asp Leu Lys Val Pro Arg Met Glu
130 135 140
Glu Lys Glu Ala Leu Val Pro Ile Gln Lys Ala Thr Asp Ser Phe His
145 150 155 160
Thr Glu Leu His Pro Arg Val Ala Phe Trp Ile Ile Lys Leu Pro Arg
165 170 175
Arg Arg Ser His Gln Asp Ala Leu Glu Gly Gly His Trp Leu Ser Glu
180 185 190
Lys Arg His Arg Leu Gln Ala Ile Arg Asp Gly Leu Arg Lys Gly Thr
195 200 205
His Lys Asp Val Leu Glu Glu Gly Thr Glu Ser Ser Ser His Ser Arg
210 215 220
Leu Ser Pro Arg Lys Thr His Leu Leu Tyr Ile Leu Arg Pro Ser Arg
225 230 235 240
Gln Leu

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<210> 5
 <211> 733
 <212> DNA
 <213> Homo sapien

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<400> 5
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agagagctcc ttgggtctca caggcctcca gagcctactc caaggcttca gccgactttt 180
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ctccgagaat gtggtggcat ccattcaacc agcggagggg agcttcgagg gtgatttgaa 420
ggtacccagg atggaggaga aggaggccct ggtaccatc cagaaggcca cggacagctt 480
ccacacagaa ctccatcccc ggggtggcctt ctggatcatt aagctgccac ggcggagggtc 540
ccaccaggat gccctggagg gcggccactg gctcagcgag aagcgacacc gcctgcaggc 600
catccgggat ggactccgca aggggaccca caaggacgtc ctagaagagg ggaccgagag 660
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tcggcagctg tag 733

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<210> 6
 <211> 242
 <212> PRT
 <213> Homo sapien

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<400> 6
Met Gly Glu Ala Ser Pro Pro Ala Pro Ala Arg Arg His Leu Leu Val
1 5 10 15
Leu Leu Leu Leu Leu Ser Thr Leu Val Ile Pro Ser Thr Ala Ala Pro
20 25 30
Ile His Asp Ala Asp Ala Gln Glu Ser Ser Leu Gly Leu Thr Gly Leu
35 40 45
Gln Ser Leu Leu Gln Gly Phe Ser Arg Leu Phe Leu Lys Gly Asn Leu
50 55 60
Leu Arg Gly Ile Asp Ser Leu Phe Ser Ala Pro Met Asp Phe Arg Gly
65 70 75 80
Leu Pro Gly Asn Tyr His Lys Glu Glu Asn Gln Glu His Gln Leu Gly
85 90 95
Asn Asn Thr Leu Ser Ser His Leu Gln Ile Asp Lys Met Thr Asp Asn
100 105 110
Lys Thr Gly Glu Val Leu Ile Ser Glu Asn Val Val Ala Ser Ile Gln
115 120 125
Pro Ala Glu Gly Ser Phe Glu Gly Asp Leu Lys Val Pro Arg Met Glu
130 135 140
Glu Lys Glu Ala Leu Val Pro Ile Gln Lys Ala Thr Asp Ser Phe His

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145      150      155      160
Thr Glu Leu His Pro Arg Val Ala Phe Trp Ile Ile Lys Leu Pro Arg
      165      170      175
Arg Arg Ser His Gln Asp Ala Leu Glu Gly Gly His Trp Leu Ser Glu
      180      185      190
Lys Arg His Arg Leu Gln Ala Ile Arg Asp Gly Leu Arg Lys Gly Thr
      195      200      205
His Lys Asp Val Leu Glu Glu Gly Thr Glu Ser Ser Ser His Ser Arg
      210      215      220
Leu Ser Pro Arg Lys Thr His Leu Leu Tyr Ile Leu Arg Pro Ser Arg
      225      230      235      240
Gln Leu

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<210> 7
 <211> 733
 <212> DNA
 <213> Homo sapien

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<400> 7
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cctgaaagggt aacctgcttc ggggcataga cagcttattc tctgccccca tggacttccg 240
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cctctccagc cacctccaga tcgacaagat gaccgacaac aagacaggag aggtgctgat 360
ctccgagaat gtggtggcat ccattcaacc agcggagggg agcttcgagg gtgatttgaa 420
ggtacccagg atggaggaga aggaggccct ggtacccatc cagaaggcca cggacagctt 480
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catccgggat ggactccgca aggggaccca caaggacgtc ctagaagagg ggaccgagag 660
ctcctcccac tccaggctgt cccccgaaa gaccactta ctgtacatcc tcaggccctc 720
tcggcagctg tag 733

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<210> 8
 <211> 242
 <212> PRT
 <213> Homo sapien

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<400> 8
Met Gly Glu Ala Ser Pro Pro Ala Pro Ala Arg Arg His Leu Leu Val
1      5      10      15
Leu Leu Leu Leu Ser Thr Leu Val Ile Pro Ser Thr Ala Ala Pro
      20      25      30
Ile His Asp Ala Asp Ala Gln Glu Ser Ser Leu Gly Leu Thr Gly Leu
      35      40      45
Gln Ser Leu Leu Gln Gly Phe Ser Arg Leu Phe Leu Lys Gly Asn Leu
      50      55      60
Leu Arg Gly Ile Asp Ser Leu Phe Ser Ala Pro Met Asp Phe Arg Gly
      65      70      75      80
Leu Pro Gly Asn Tyr His Lys Glu Glu Asn Gln Glu His Gln Leu Gly
      85      90      95
Asn Asn Thr Leu Ser Ser His Leu Gln Ile Asp Lys Met Thr Asp Asn
      100      105      110
Lys Thr Gly Glu Val Leu Ile Ser Glu Asn Val Val Ala Ser Ile Gln
      115      120      125
Pro Ala Glu Gly Ser Phe Glu Gly Asp Leu Lys Val Pro Arg Met Glu
      130      135      140
Glu Lys Glu Ala Leu Val Pro Ile Gln Lys Ala Thr Asp Ser Phe His
145      150      155      160
Thr Glu Leu His Pro Arg Val Ala Phe Trp Ile Ile Lys Leu Pro Arg
      165      170      175

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Arg Arg Ser His Gln Asp Ala Leu Glu Gly Gly His Trp Leu Ser Glu
 180 185 190
 Lys Arg His Arg Leu Gln Ala Ile Arg Asp Gly Leu Arg Lys Gly Thr
 195 200 205
 His Lys Asp Val Leu Glu Glu Gly Thr Glu Ser Ser Ser His Ser Arg
 210 215 220
 Leu Ser Pro Arg Lys Thr His Leu Leu Tyr Ile Leu Arg Pro Ser Arg
 225 230 235 240
 Gln Leu

<210> 9
 <211> 733
 <212> DNA
 <213> Homo sapien

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 agagagctcc ttgggtctca caggcctcca gagcctactc caaggcttca gccgactttt 180
 cctgaaaggt aacctgcttc ggggcataga cagcttattc tctgccccca tggacttccg 240
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 tcggcagctg tag 733

<210> 10
 <211> 242
 <212> PRT
 <213> Homo sapien

<400> 10
 Met Gly Glu Ala Ser Pro Pro Ala Pro Ala Arg Arg His Leu Leu Val
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 Leu Leu Leu Leu Leu Ser Thr Leu Val Ile Pro Ser Thr Ala Ala Pro
 20 25 30
 Ile His Asp Ala Asp Ala Gln Glu Ser Ser Leu Gly Leu Thr Gly Leu
 35 40 45
 Gln Ser Leu Leu Gln Gly Phe Ser Arg Leu Phe Leu Lys Gly Asn Leu
 50 55 60
 Leu Arg Gly Ile Asp Ser Leu Phe Ser Ala Pro Met Asp Phe Arg Gly
 65 70 75 80
 Leu Pro Gly Asn Tyr His Lys Glu Glu Asn Gln Glu His Gln Leu Gly
 85 90 95
 Asn Asn Thr Leu Ser Ser His Leu Gln Ile Asp Lys Met Thr Asp Asn
 100 105 110
 Lys Thr Gly Glu Val Leu Ile Ser Glu Asn Val Val Ala Ser Ile Gln
 115 120 125
 Pro Ala Glu Gly Ser Phe Glu Gly Asp Leu Lys Val Pro Arg Met Glu
 130 135 140
 Glu Lys Glu Ala Leu Val Pro Ile Gln Lys Ala Thr Asp Ser Phe His
 145 150 155 160
 Thr Glu Leu His Pro Arg Val Ala Phe Trp Ile Ile Lys Leu Pro Arg
 165 170 175
 Arg Arg Ser His Gln Asp Ala Leu Glu Gly Ser His Trp Leu Ser Glu
 180 185 190
 Lys Arg His Arg Leu Gln Ala Ile Arg Asp Gly Leu Arg Lys Gly Thr

SEQLIST.TXT

195 200 205
 His Lys Asp Val Leu Lys Glu Thr Glu Ser Ser Ser His Ser Arg
 210 215 220
 Leu Ser Pro Arg Lys Thr His Leu Leu Tyr Ile Leu Arg Pro Ser Arg
 225 230 235 240
 Gln Leu

<210> 11
 <211> 733
 <212> DNA
 <213> Homo sapien

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 cctgaaagggt aacctgcttc ggggcataga cagcttattc tctgccccca tggacttccg 240
 gggcctccct gggaactacc acaaagagga gaaccaggag caccagctgg ggaacaacac 300
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 ctccgagaat gtggtggcat ccattcaacc agcggagggg agcttcgagg gtgatttgaa 420
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 catccgggat ggactccgca aggggaccca caaggacgtc ctagaagagg ggaccgagag 660
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<210> 12
 <211> 242
 <212> PRT
 <213> Homo sapien

<400> 12
 Met Gly Glu Ala Ser Pro Pro Ala Pro Ala Arg Arg His Leu Leu Val
 1 5 10 15
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 20 25 30
 Ile His Asp Ala Asp Ala Gln Glu Ser Ser Leu Gly Leu Thr Gly Leu
 35 40 45
 Gln Ser Leu Leu Gln Gly Phe Ser Arg Leu Phe Leu Lys Gly Asn Leu
 50 55 60
 Leu Arg Gly Ile Asp Ser Leu Phe Ser Ala Pro Met Asp Phe Arg Gly
 65 70 75 80
 Leu Pro Gly Asn Tyr His Lys Glu Glu Asn Gln Glu His Gln Leu Gly
 85 90 95
 Asn Asn Thr Leu Ser Ser His Leu Gln Ile Asp Lys Met Thr Asp Asn
 100 105 110
 Lys Thr Gly Glu Val Leu Ile Ser Glu Asn Val Val Ala Ser Ile Gln
 115 120 125
 Pro Ala Glu Gly Ser Phe Glu Gly Asp Leu Lys Val Pro Arg Met Glu
 130 135 140
 Glu Lys Glu Ala Leu Val Pro Ile Gln Lys Ala Thr Asp Ser Phe His
 145 150 155 160
 Thr Glu Leu His Pro Arg Val Ala Phe Trp Ile Ile Lys Leu Pro Arg
 165 170 175
 Arg Arg Ser His Gln Asp Ala Leu Glu Gly Gly His Trp Leu Ser Glu
 180 185 190
 Lys Arg His Arg Leu Gln Ala Ile Arg Asp Gly Leu Arg Lys Gly Thr
 195 200 205
 His Lys Asp Val Leu Glu Glu Gly Thr Glu Ser Ser Ser His Ser Arg
 210 215 220

SEQLIST.TXT

Leu Ser Pro Arg Lys Thr His Leu Leu Tyr Ile Leu Arg Pro Ser Arg
 225 230 235 240
 Gln Leu

<210> 13
 <211> 640
 <212> DNA
 <213> Homo sapien
 <400> 13
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 acccatccag aaggccacgg acagcttcca cacagaactc catccccggg tggccttctg 420
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 ccacttactg tacatcctca ggccctctcg gcagctgtag 640

<210> 14
 <211> 211
 <212> PRT
 <213> Homo sapien
 <400> 14
 Met Gly Glu Ala Ser Pro Pro Ala Pro Ala Arg Arg His Leu Leu Val
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 Leu Leu Leu Leu Ser Thr Leu Val Ile Pro Ser Ala Ala Ala Pro
 20 25 30
 Ile His Asp Ala Asp Ala Gln Glu Ser Ser Leu Gly Leu Thr Gly Leu
 35 40 45
 Gln Ser Leu Leu Gln Gly Phe Ser Arg Leu Phe Leu Lys Gly Asn Leu
 50 55 60
 Leu Arg Gly Ile Asp Ser Leu Phe Ser Ala Pro Met Asp Phe Arg Gly
 65 70 75 80
 Leu Pro Gly Asn Tyr His Lys Glu Glu Asn Gln Glu His Gln Leu Gly
 85 90 95
 Asn Asn Thr Leu Ser Ser His Leu Gln Ile Asp Lys Val Pro Arg Met
 100 105 110
 Glu Glu Lys Glu Ala Leu Val Pro Ile Gln Lys Ala Thr Asp Ser Phe
 115 120 125
 His Thr Glu Leu His Pro Arg Val Ala Phe Trp Ile Ile Lys Leu Pro
 130 135 140
 Arg Arg Arg Ser His Gln Asp Ala Leu Glu Gly Gly His Trp Leu Ser
 145 150 155 160
 Glu Lys Arg His Arg Leu Gln Ala Ile Arg Asp Gly Leu Arg Lys Gly
 165 170 175
 Thr His Lys Asp Val Leu Glu Glu Gly Thr Glu Ser Ser Ser His Ser
 180 185 190
 Arg Leu Ser Pro Arg Lys Thr His Leu Leu Tyr Ile Leu Arg Pro Ser
 195 200 205
 Arg Gln Leu
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<210> 15
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 <212> DNA
 <213> Homo sapien

SEQLIST.TXT

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<400> 15
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cctctccagc cacctccaga tcgacaaggt acccaggatg gaggagaagg aggccctggg 360
acccatccag aaggccacgg acagcttcca cacagaactc catccccggg tggccttctg 420
gatcattaag ctgccacggc ggaggtccca ccaggatgcc ctggaggggc gccactggct 480
cagcgagaag cgacaccgcc tgcaggccat ccgggatgga ctccgcaagg ggacccacaa 540
ggacgtccta gaagaggaga ccgagagctc ctcccactcc aggcgtgtccc cccgaaagac 600
ccacttactg tacatctca ggcctctctg gcagctgtag 640

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<210> 16
<211> 211
<212> PRT
<213> Homo sapien

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<400> 16
Met Gly Glu Ala Ser Pro Pro Ala Pro Ala Arg Arg His Leu Leu Val
1      5      10      15
Leu Leu Leu Leu Leu Ser Thr Leu Val Ile Pro Ser Ala Ala Ala Pro
20     25     30
Ile His Asp Ala Asp Ala Gln Glu Ser Ser Leu Gly Leu Thr Gly Leu
35     40     45
Gln Ser Leu Leu Gln Gly Phe Ser Arg Leu Phe Leu Lys Gly Asn Leu
50     55     60
Leu Arg Gly Ile Asp Ser Leu Phe Ser Ala Pro Met Asp Phe Arg Gly
65     70     75     80
Leu Pro Gly Asn Tyr His Lys Glu Glu Asn Gln Glu His Gln Leu Gly
85     90     95
Asn Asn Thr Leu Ser Ser His Leu Gln Ile Asp Lys Val Pro Arg Met
100    105    110
Glu Glu Lys Glu Ala Leu Val Pro Ile Gln Lys Ala Thr Asp Ser Phe
115    120    125
His Thr Glu Leu His Pro Arg Val Ala Phe Trp Ile Ile Lys Leu Pro
130    135    140
Arg Arg Arg Ser His Gln Asp Ala Leu Glu Gly Gly His Trp Leu Ser
145    150    155    160
Glu Lys Arg His Arg Leu Gln Ala Ile Arg Asp Gly Leu Arg Lys Gly
165    170    175
Thr His Lys Asp Val Leu Glu Glu Glu Thr Glu Ser Ser Ser His Ser
180    185    190
Arg Leu Ser Pro Arg Lys Thr His Leu Leu Tyr Ile Leu Arg Pro Ser
195    200    205
Arg Gln Leu
210

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<210> 17
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<212> DNA
<213> Homo sapien

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<400> 17
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gcaggccatc cgggatggac tccgcaaggg gacccacaag gacgtcctaa aagaggggac 420
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<210> 18
<211> 164
<212> PRT
<213> Homo sapien

<400> 18

Met	Gly	Glu	Ala	Ser	Pro	Pro	Ala	Pro	Ala	Arg	Arg	His	Leu	Leu	Val
1				5				10					15		
Leu	Leu	Leu	Leu	Leu	Ser	Thr	Leu	Val	Ile	Pro	Ser	Ala	Ala	Ala	Pro
			20				25					30			
Ile	His	Asp	Ala	Asp	Ala	Gln	Glu	Ser	Ser	Leu	Gly	Leu	Thr	Gly	Leu
		35				40					45				
Gln	Ser	Leu	Leu	Gln	Gly	Phe	Ser	Arg	Leu	Phe	Leu	Lys	Val	Pro	Arg
	50				55					60					
Met	Glu	Glu	Lys	Glu	Ala	Leu	Val	Pro	Ile	Gln	Lys	Ala	Thr	Asp	Ser
65				70				75					80		
Phe	His	Thr	Glu	Leu	His	Pro	Arg	Val	Ala	Phe	Trp	Ile	Ile	Lys	Leu
			85					90					95		
Pro	Arg	Arg	Arg	Ser	His	Gln	Asp	Ala	Leu	Glu	Gly	Ser	His	Trp	Leu
			100					105					110		
Ser	Glu	Lys	Arg	His	Arg	Leu	Gln	Ala	Ile	Arg	Asp	Gly	Leu	Arg	Lys
		115					120					125			
Gly	Thr	His	Lys	Asp	Val	Leu	Lys	Glu	Gly	Thr	Glu	Ser	Ser	Ser	His
	130					135					140				
Ser	Arg	Leu	Ser	Pro	Arg	Lys	Thr	His	Leu	Leu	Tyr	Ile	Leu	Arg	Pro
145					150					155					160
Ser	Arg	Gln	Leu												

<210> 19
<211> 499
<212> DNA
<213> Homo sapien

<400> 19

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agagagctcc	ttgggtctca	caggcctcca	gagcctactc	caaggcttca	gccgactttt	180
cctgaaagta	cccaggatgg	aggagaagga	ggcccttgta	cccatccaga	aggccacgga	240
cagcttccac	acagaactcc	atccccgggt	ggccttctgg	atcattaagc	tgccacggcg	300
gaggtcccac	caggatgccc	tggagggcag	ccactggctc	agcgagaagc	gacaccgcct	360
gcaggccatc	cgggatggac	tccgcaaggg	gacccacaag	gacgtcctaa	aagaggggac	420
cgagagctcc	tcccactcca	ggctgtcccc	ccgaaagacc	cacttactgt	acatcctcag	480
gccctctcgg	cagctgtag					499

<210> 20
<211> 164
<212> PRT
<213> Homo sapien

<400> 20

Met	Gly	Glu	Ala	Ser	Pro	Pro	Ala	Pro	Ala	Arg	Arg	His	Leu	Leu	Val
1				5				10					15		
Leu	Leu	Leu	Leu	Leu	Ser	Thr	Leu	Val	Ile	Pro	Ser	Ala	Ala	Ala	Pro
			20				25					30			
Ile	His	Asp	Ala	Asp	Ala	Gln	Glu	Ser	Ser	Leu	Gly	Leu	Thr	Gly	Leu
		35				40					45				
Gln	Ser	Leu	Leu	Gln	Gly	Phe	Ser	Arg	Leu	Phe	Leu	Lys	Val	Pro	Arg
	50				55					60					
Met	Glu	Glu	Lys	Glu	Ala	Leu	Val	Pro	Ile	Gln	Lys	Ala	Thr	Asp	Ser
65				70				75					80		

SEQLIST.TXT

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Phe His Thr Glu Leu His Pro Arg Val Ala Phe Trp Ile Ile Lys Leu
      85      90
Pro Arg Arg Arg Ser His Gln Asp Ala Leu Glu Gly Ser His Trp Leu
      100      105      110
Ser Glu Lys Arg His Arg Leu Gln Ala Ile Arg Asp Gly Leu Arg Lys
      115      120      125
Gly Thr His Lys Asp Val Leu Lys Glu Gly Thr Glu Ser Ser Ser His
      130      135      140
Ser Arg Leu Ser Pro Arg Lys Thr His Leu Leu Tyr Ile Leu Arg Pro
      145      150      155      160
Ser Arg Gln Leu

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<210> 21
<211> 499
<212> DNA
<213> Homo sapien

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<400> 21
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agagagctcc ttgggtctca caggcctcca gagcctactc caaggcttca gccgactttt 180
cctgaaagta cccaggatgg aggagaagga ggccctggta cccatccaga aggccacgga 240
cagcttccac acagaactcc atccccgggt ggcttctctg atcattaagc tgccacggcg 300
gaggtcccac caggatgccc tggaggcag ccactggctc agcgagaagc gacaccgcct 360
gcaggccatc cgggatggac tccgcaaggg gaccacaag gacgtcctag aagaggggac 420
cgagagctcc tccactcca ggctgtcccc ccgaaagacc cacttactgt acatcctcag 480
gccctctcgg cagctgtag                                     499

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<210> 22
<211> 164
<212> PRT
<213> Homo sapien

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<400> 22
Met Gly Glu Ala Ser Pro Pro Ala Pro Ala Arg Arg His Leu Leu Val
 1      5      10      15
Leu Leu Leu Leu Leu Ser Thr Leu Val Ile Pro Ser Ala Ala Ala Pro
      20      25      30
Ile His Asp Ala Asp Ala Gln Glu Ser Ser Leu Gly Leu Thr Gly Leu
      35      40      45
Gln Ser Leu Leu Gln Gly Phe Ser Arg Leu Phe Leu Lys Val Pro Arg
      50      55      60
Met Glu Glu Lys Glu Ala Leu Val Pro Ile Gln Lys Ala Thr Asp Ser
      65      70      75      80
Phe His Thr Glu Leu His Pro Arg Val Ala Phe Trp Ile Ile Lys Leu
      85      90      95
Pro Arg Arg Arg Ser His Gln Asp Ala Leu Glu Gly Ser His Trp Leu
      100      105      110
Ser Glu Lys Arg His Arg Leu Gln Ala Ile Arg Asp Gly Leu Arg Lys
      115      120      125
Gly Thr His Lys Asp Val Leu Glu Glu Gly Thr Glu Ser Ser Ser His
      130      135      140
Ser Arg Leu Ser Pro Arg Lys Thr His Leu Leu Tyr Ile Leu Arg Pro
      145      150      155      160
Ser Arg Gln Leu

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<210> 23
<211> 499
<212> DNA

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SEQLIST.TXT

<213> Homo sapien

<400> 23

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agagagctcc ttgggtctca caggcctcca gagcctactc caaggcttca gccgactttt 180
cctgaaagta cccaggatgg aggagaagga ggccctggta cccatccaga aggccacgga 240
cagcttccac acagaactcc atccccgggt ggccttctgg atcattaagc tgccacggcg 300
gaggtcccac caggatgccc tggagggcag ccactggctc agcgagaagc gacaccgcct 360
gcaggccatc cgggatggac tccgcaaggg gaccacaaag gacgtcctaa aagaggggac 420
cgagagctcc tcccactcca ggctgtcccc ccgaaagacc cacttactgt acatcctcag 480
gccctctcgg cagctgtag                                     499

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<210> 24

<211> 164

<212> PRT

<213> Homo sapien

<400> 24

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Met Gly Glu Ala Ser Pro Pro Ala Pro Ala Arg Arg His Leu Leu Val
1      5      10      15
Leu Leu Leu Leu Leu Ser Thr Leu Val Ile Pro Ser Ala Ala Ala Pro
20     25     30
Ile His Asp Ala Asp Ala Gln Glu Ser Ser Leu Gly Leu Thr Gly Leu
35     40     45
Gln Ser Leu Leu Gln Gly Phe Ser Arg Leu Phe Leu Lys Val Pro Arg
50     55     60
Met Glu Glu Lys Glu Ala Leu Val Pro Ile Gln Lys Ala Thr Asp Ser
65     70     75     80
Phe His Thr Glu Leu His Pro Arg Val Ala Phe Trp Ile Ile Lys Leu
85     90     95
Pro Arg Arg Arg Ser His Gln Asp Ala Leu Glu Gly Ser His Trp Leu
100    105    110
Ser Glu Lys Arg His Arg Leu Gln Ala Ile Arg Asp Gly Leu Arg Lys
115    120    125
Gly Thr His Lys Asp Val Leu Lys Glu Gly Thr Glu Ser Ser Ser His
130    135    140
Ser Arg Leu Ser Pro Arg Lys Thr His Leu Leu Tyr Ile Leu Arg Pro
145    150    155    160
Ser Arg Gln Leu

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<210> 25

<211> 499

<212> DNA

<213> Homo sapien

<400> 25

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agagagctcc ttgggtctca caggcctcca gagcctactc caaggcttca gccgactttt 180
cctgaaagta cccaggatgg aggagaagga ggccctggta cccatccaga aggccacgga 240
cagcttccac acagaactcc atccccgggt ggccttctgg atcattaagc tgccacggcg 300
gaggtcccac caggatgccc tggagggcag ccactggctc agcgagaagc gacaccgcct 360
gcaggccatc cgggatggac tccgcaaggg gaccacaaag gacgtcctaa aagaggggac 420
cgagagctcc tcccactcca ggctgtcccc ccgaaagacc cacttactgt acatcctcag 480
gccctctcgg cagctgtag                                     499

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<210> 26

<211> 164

<212> PRT

<213> Homo sapien

SEQLIST.TXT

<400> 26

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Met Gly Glu Ala Ser Pro Pro Ala Pro Ala Arg Arg His Leu Leu Val
 1      5      10      15
Leu Leu Leu Leu Leu Ser Thr Leu Val Ile Pro Ser Thr Ala Ala Pro
 20      25      30
Ile His Asp Ala Asp Ala Gln Glu Ser Ser Leu Gly Leu Thr Gly Leu
 35      40      45
Gln Ser Leu Leu Gln Gly Phe Ser Arg Leu Phe Leu Lys Val Pro Arg
 50      55      60
Met Glu Glu Lys Glu Ala Leu Val Pro Ile Gln Lys Ala Thr Asp Ser
 65      70      75      80
Phe His Thr Glu Leu His Pro Arg Val Ala Phe Trp Ile Ile Lys Leu
 85      90      95
Pro Arg Arg Arg Ser His Gln Asp Ala Leu Glu Gly Ser His Trp Leu
100      105      110
Ser Glu Lys Arg His Arg Leu Gln Ala Ile Arg Asp Gly Leu Arg Lys
115      120      125
Gly Thr His Lys Asp Val Leu Lys Glu Gly Thr Glu Ser Ser Ser His
130      135      140
Ser Arg Leu Ser Pro Arg Lys Thr His Leu Leu Tyr Ile Leu Arg Pro
145      150      155      160
Ser Arg Gln Leu

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<210> 27

<211> 20

<212> DNA

<213> Homo sapien

<400> 27

atcgacaagg tacccaggat

20

<210> 28

<211> 20

<212> DNA

<213> Homo sapien

<400> 28

ttcctgaaag tacccaggat

20